

Fri Mar 2 09:30:23 2001

us-09-331-631a-5_copy_76_144.rag

GenCore version 4.5
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OM protein - protein search, using sw model

Run on: March 1, 2001, 15:47:15 ; Search time 210.42 Seconds

(Without alignments)
11.213 Million cell updates/sec

Title: US-09-331-631A-5_COPY_76_144
Perfect score: 381
Sequence: 1 NRORDDQQQYEQCOKRCOR.....EEQREDEKYEERMKESGN 69

Scoring table:

BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 268485 seqs, 34193795 residues

Total number of hits satisfying chosen parameters: 268485

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%
Listing first 45 summaries

Database :

A-Geneseq_36:*

1: /SIDS1/gcgdata/geneseq/AA1980.DAT:*

2: /SIDS1/gcgdata/geneseq/AA1981.DAT:*

3: /SIDS1/gcgdata/geneseq/AA1982.DAT:*

4: /SIDS1/gcgdata/geneseq/AA1983.DAT:*

5: /SIDS1/gcgdata/geneseq/AA1984.DAT:*

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9: /SIDS1/gcgdata/geneseq/AA1988.DAT:*

10: /SIDS1/gcgdata/geneseq/AA1989.DAT:*

11: /SIDS1/gcgdata/geneseq/AA1990.DAT:*

12: /SIDS1/gcgdata/geneseq/AA1991.DAT:*

13: /SIDS1/gcgdata/geneseq/AA1992.DAT:*

14: /SIDS1/gcgdata/geneseq/AA1993.DAT:*

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16: /SIDS1/gcgdata/geneseq/AA1995.DAT:*

17: /SIDS1/gcgdata/geneseq/AA1996.DAT:*

18: /SIDS1/gcgdata/geneseq/AA1997.DAT:*

19: /SIDS1/gcgdata/geneseq/AA1998.DAT:*

20: /SIDS1/gcgdata/geneseq/AA1999.DAT:*

21: /SIDS1/gcgdata/geneseq/AA2000.DAT:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result	Score	Query Match	Length	DB ID	Description
1	381	100.0	625	19	W62830
2	363	95.3	666	19	W62828
3	359	94.2	666	19	W62829
4	173	45.4	525	13	W62831
5	173	45.4	566	13	R20181
6	152.5	40.0	590	19	W62832
7	103	27.0	1898	20	W62832
8	102	26.8	1162	21	Y50795
9	97.5	25.6	1135	21	Y58500
10	97.5	25.6	1233	21	Y68784
11	97.5	25.6	1239	20	Y55954
12	97	25.5	482	20	Y55931
					Renal cancer assoc

13	97	25.5	1297	20	Y32154
14	96	25.2	1360	20	Y32154
15	95.5	25.1	1326	20	Y32154
16	95	24.9	444	20	Y32154
17	95	24.9	524	20	Y32154
18	94.5	24.8	611	20	Y32154
19	93	24.4	2023	21	Y54311
20	93	24.4	2074	21	Y54311
21	90.5	23.8	409	20	W90342
22	90.5	23.8	489	20	W90341
23	89	23.4	346	20	W90115
24	89	23.4	373	20	W90115
25	89	23.1	593	19	W62835
26	88	23.1	1197	21	Y57445
27	88	23.1	1658	21	Y57445
28	86.5	22.7	562	16	R70491
29	85.5	22.4	740	13	R27530
30	85.5	22.4	740	16	R68838
31	85	22.3	910	20	Y22191
32	85	22.3	1214	21	Y57444
33	85	22.3	1715	21	Y57449
34	84.5	22.2	1299	21	Y56633
35	84	22.0	200	18	W55301
36	84	22.0	200	18	W20685
37	84	22.0	529	18	W50444
38	84	22.0	1382	18	W31867
39	83.5	21.9	303	15	R60054
40	83.5	21.9	356	20	Y20119
41	83.5	21.9	326	20	Y20118
42	82.5	21.7	432	20	W93954
43	82	21.5	301	8	P70867
44	82	21.5	648	20	Y32157
45	82	21.5	1144	20	Y32154

ALIGNMENTS

RESULT 1	W62830	standard: Protein; 625 AA.
ID	W62830	
AC	W62830	
XX		
DT	27-OCT-1998	(first entry)
DE	Macadamia integrifolia antimicrobial protein.	
KM	antimicrobial protein; infestation; control.	
XX		
OS	Macadamia integrifolia.	
XX		
EH	Key	Location/Qualifiers
FT	Peptide	1..28
FT	Protein	/note="signal peptide"
FT		29..666
XX		/note="mature protein"
PN	W09827805-A1.	
XX		
PD	02-JUL-1998.	
XX		
EP	22-DEC-1997;	97WO-AU00874.
XX		
PR	20-DEC-1996;	96AU-0004275.
XX		
PA	(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.	
XX		
PI	Bower NI, Coulter KC, Green JL, Manners JM, Marcus JP;	
DR	WPI: 1998-372729/32.	
XX	N-PSDB; V42316.	
XX		

Novel anti-microbial protein from e.g. Macadamia integrifolia -
useful for controlling microbial infestations of plants or mammals
Claim 1: Page 43-45; 96pp; English.

The sequence is that of an antimicrobial protein which can
be used to control microbial infestations in plants and mammalian
animals.

Sequence 625 AA:

Query Match 100.0%; Score 381; DB 19; Length 625;
Best Local Similarity 100.0%; Pred. No. 1.8e-30;
Matches 69; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

1 NRQRPDPOOYEGCCORRCORRETEPRHMOICQRCERREYKRRKQKRYEEOOREDEEKY 60
|||||
b 76 nrqrdpqgqyegqcrqrreterprimqicqrcerryekrkxqkryeeqredeeky 135
|||||
y 61 EERMKEGDN 69
|||||
db 136 eermekegdn 144

RESULT 2

W62828 standard; Protein; 666 AA.

W62828; 27-OCT-1998 (first entry)

Macadamia integrifolia antimicrobial protein.
antimicrobial protein; infestation; control.
Macadamia integrifolia.

Key Location/Qualifiers
Peptide 1..28
FT /note="signal peptide"
FT 29..666
FT Protein /note="mature protein"

W09827805-A1.

02-JUL-1998.

22-DEC-1997; 97WO-AU00874.

20-DEC-1996; 96AU-0004275.

(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP;

WPI; 1998-377279/32.

N-PSDB; V42310.

Novel anti-microbial protein from e.g. Macadamia integrifolia -
useful for controlling microbial infestations of plants or mammals

Claim 1: Page 34-36; 96pp; English.

The sequence is that of an antimicrobial protein which can
be used to control microbial infestations in plants and mammalian
animals.

Sequence 666 AA;

Query Match

95 3%, Score 363; DB 19; Length 666;

Best Local Similarity 95.7%; Pred. No. 1.2e-28;
Matches 66; Conservative 0; Mismatches 3; Indels 0; Gaps 0.

QY 1 NRQRPDPOOYEGCCORRCORRETEPRHMOICQRCERREYKRRKQKRYEEOOREDEEKY 60
|||||

Db 117 nrqrdpqgqyegqcrqrreterprimqicqrcerryekrkxqkryeeqredeeky 176
|||||

QY 61 EERMKEGDN 69
|||||

Db 177 eermekegdn 185

RESULT 3

W62829 standard; Protein; 666 AA.

W62829; 27-OCT-1998 (first entry)

Macadamia integrifolia antimicrobial protein.
antimicrobial protein; infestation; control.
Macadamia integrifolia.

Key Location/Qualifiers
Peptide 1..28
FT /note="signal peptide"
FT 29..666
FT Protein /note="mature protein"

W09827805-A1.

02-JUL-1998.

22-DEC-1997; 97WO-AU00874.

20-DEC-1996; 96AU-0004275.

(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP;

WPI; 1998-377279/32.

N-PSDB; V42311.

Novel anti-microbial protein from e.g. Macadamia integrifolia -
useful for controlling microbial infestations of plants or mammals

Claim 1: Page 39-41; 96pp; English.

The sequence is that of an antimicrobial protein which can
be used to control microbial infestations in plants and mammalian
animals.

Sequence 666 AA;

Query Match 94.2%; Score 359; DB 19;
Best Local Similarity 94.2%; Pred. No. 2.9e-27;
Matches 65; Conservative 1; Mismatches 3

QY 1 NRQRPDPOOYEGCCORRCORRETEPRHMOICQRCERREYKRRKQKRYEEOOREDEEKY 60
|||||

Db 117 nrqrdpqgqyegqcrqrreterprimqicqrcerryekrkxqkryeeqredeeky 176
|||||

QY 61 EERMKEGDN 69
|||||

Db 177 eermekegdn 185

RESULT 4

```

W62831
ID W62831 standard; Protein; 525 AA.
XX AC
AC W62831;
XX DT
DT 27-OCT-1998 (first entry)
XX DE
DE Theobroma cacao antimicrobial protein.
XX KM
KM antimicrobial protein; infestation; control.
XX OS
OS Theobroma cacao.
XX PN
PN WO9827805-A1.
XX PD
PD 02-JUL-1998.
XX PF
PF 22-DEC-1997; 97WO-AU00874.
XX PR
PR 20-DEC-1996; 96AU-0004275.
XX PA
PA (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
XX PI
PI Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP;
XX DR
DR WPI; 1998-377279/32.
XX PT
PT Novel anti-microbial protein from e.g. Macadamia integrifolia -
XX PS
PS useful for controlling microbial infestations of plants or mammals
XX CC
CC Claim 1; Page 47-49; 96pp; English.
XX CC
CC The sequence is that of an antimicrobial protein which can
XX CC
CC be used to control microbial infestations in plants and mammalian
XX CC
CC animals.
SQ Sequence 525 AA:

Query Match 45.4%; Score 173; DB 19; Length 525;
Best Local Similarity 32.6%; Pred. No. 6.7e-10;
Matches 31; Conservative 17; Mismatches 15; Indels 32; Gaps 1.

QY 3 QRPDOOVYEQCCKRCGRREPERHMQICORCERYEKEKRRKO----- 46
DB 35 erdprqyegcqqrceaeleeregegcqrceteykqgrqqeeelrqyggcggrgqe 94
   :|||:|||||||:|:-|||:|:||||| |:::|:|:|
OY 47 -----KRYEEQQREDEEKYEERNK 65
   ::|||:|:| |
DB 95 qggqgreqgcqqrckwecqykegerghenyhnkk 129

RESULT 5
R20181 ID R20181 standard; Protein; 566 AA.
XX AC
AC R20181;
XX DT
DT 16-APR-1992 (first entry)
XX DE
DE Sequence encoded by 67 kD T. cacao protein cDNA.
XX KM
KM Cocoa; flavour; vicillin; seed storage protein.
XX OS
OS Theobroma cacao.
XX PN
PN WO9119801-A.
XX PD
PD 26-DEC-1991.
XX PF
PF 07-JUN-1991; 91WO-GH00914.
XX PR
PR 11-JUN-1990; 90GB-0013016.
```

[illegible]

[illegible]

FT	Modified-site	/note=	"potential phosphorylation site"
FT	Modified-site	/note=	"potential phosphorylation site"
FT	Modified-site	543	"potential phosphorylation site"
FT	Modified-site	/note=	"potential phosphorylation site"
FT	Modified-site	550	"potential phosphorylation site"
FT	Modified-site	/note=	"potential phosphorylation site"
FT	Modified-site	554	"potential phosphorylation site"
FT	Modified-site	/note=	"potential phosphorylation site"
FT	Modified-site	570	"potential phosphorylation site"
FT	Modified-site	/note=	"potential glycosylation site"
FT	Modified-site	572	"potential phosphorylation site"
FT	Modified-site	/note=	"potential phosphorylation site"
FT	Modified-site	624	"potential phosphorylation site"
FT	Modified-site	/note=	"potential phosphorylation site"
FT	Modified-site	625	"potential phosphorylation site"
FT	Modified-site	/note=	"potential phosphorylation site"
FT	Modified-site	632	"potential phosphorylation site"
FT	Modified-site	/note=	"potential phosphorylation site"
FT	Modified-site	681	"potential phosphorylation site"
FT	Modified-site	/note=	"potential phosphorylation site"
FT	Modified-site	682	"potential phosphorylation site"
FT	Modified-site	/note=	"potential phosphorylation site"
FT	Modified-site	688	"potential phosphorylation site"
FT	Modified-site	/note=	"potential phosphorylation site"
FT	Modified-site	689	"potential phosphorylation site"
FT	Modified-site	/note=	"potential phosphorylation site"
FT	Modified-site	706	"potential phosphorylation site"
FT	Modified-site	/note=	"potential phosphorylation site"
FT	Modified-site	718	"potential phosphorylation site"
FT	Modified-site	/note=	"potential glycosylation site"
FT	Modified-site	720	"potential phosphorylation site"
FT	Modified-site	/note=	"potential phosphorylation site"
FT	Modified-site	726	"potential phosphorylation site"
FT	Modified-site	/note=	"potential phosphorylation site"
FT	Modified-site	811	"potential phosphorylation site"
FT	Modified-site	/note=	"potential phosphorylation site"
FT	Modified-site	815	"potential phosphorylation site"
FT	Modified-site	/note=	"potential phosphorylation site"
FT	Domain	836..1115	"potential phosphorylation site"
FT	Modified-site	/note=	"NIK1-like kinase domain"
FT	Modified-site	898	"potential phosphorylation site"
FT	Modified-site	/note=	"potential phosphorylation site"
FT	Modified-site	931	"potential phosphorylation site"
FT	Modified-site	/note=	"potential phosphorylation site"
FT	Modified-site	958	"potential phosphorylation site"
FT	Modified-site	/note=	"potential phosphorylation site"
FT	Modified-site	978	"potential phosphorylation site"
FT	Modified-site	/note=	"potential phosphorylation site"
FT	Modified-site	999	"potential phosphorylation site"
FT	Modified-site	/note=	"potential phosphorylation site"
FT	Modified-site	1012	"potential phosphorylation site"
FT	Modified-site	/note=	"potential phosphorylation site"
FT	Modified-site	1067	"potential phosphorylation site"
FT	Modified-site	/note=	"potential glycosylation site"
FT	Modified-site	1113	"potential phosphorylation site"
XX	W0200006728-A2.	/note=	"potential phosphorylation site"
XX	10-FEB-2000.		
XX	28-JUL-1999;	99W0-US17132.	
XX	28-JUL-1998;	98US-0123434.	
XX	14-SEP-1998;	98US-0152814.	
PR	14-OCT-1998;	98US-0173482.	
PR	03-NOV-1998;	98US-0106889.	
PR	19-NOV-1998;	98US-0109093.	
PR	22-DEC-1998;	98US-0113796.	
PR	12-JAN-1999;	98US-0173482.	
XX	12-JAN-1999;	99US-0229005.	

PA	(INCY)-INCYTE PHARM INC.
XX	
Pt	Hillman JL, Lal P, Tang YT, Corley NC, Guegler KJ, Baughn MR;
Pt	Patterson C, Bandman O, Au-Young J, Gorgone GA, Yue R, Azimzai Y;
Pt	Reddy R, Lu DAM, Shih LL;
XX	
DR	WPI: 2000-183125/16.
DR	N-PSDB: Z46153.
XX	
PT	New human phosphorylation effectors useful for the diagnosis, treatment
PT	and prevention of proliferative, immune and neuronal disorders -
PS	
PS	Claim 1: Page 98-100; 142pp: English.
XX	
CC	Y68769-95 and Y68797-99 represent human phosphorylation effectors (PHSP),
CC	designated PHSP1-PHSP31 (the protein sequence for PHSP28 is not given
CC	in the specification). The sequences were isolated from cDNA libraries
CC	prepared from various human tissues. The PHSP proteins are useful for
CC	the diagnosis, treatment and prevention of proliferative disorders,
CC	immune disorders and neuronal disorders. The PHSP proteins form
CC	pharmaceutical compositions which useful for treating or preventing
CC	disorders associated with decreased PHSP expression/activity. PHSP
CC	antagonists are useful for treating or preventing disorders associated
CC	with increased PHSP expression/activity.
SQ	
SQ	Sequence 1135 AA:
	Query Match 25.6%; Score 97.5; DB 21; Length 1135;
	Best Local Similarity 39.2%; Pred. No. 0.047;
	Matches 29; Conservative 17; Mismatches 17; Indels 11; Gaps 4
QY	2 RQRDPOOYEQCK--RCORRETPRHMOICQRCRRERKEKRN---QOKRYEQQR- 54
Dd	11:: :11 11 :: 1111 1 1 1 :11 11 :1111: :::111::1
Dd	395 rtkrlieqkqqrtrrleeqqrrrearrqreqqr--rreeqekrrleelerrkeeeerr 452
QY	55 --EDEKYEERMKE 66
Dd	1::1:: 11::1
Dd	453 raeeekrrveeqe 466
RESULT 10	
Y55954	
ID	Y55954 standard; Protein: 1233 AA.
AC	
AC	Y55954:
XX	
DT	18-FEB-2000 (first entry)
XX	
DE	Mouse STE20-related protein kinase NIK_m.
XX	
RW	Antirheumatic; antiarthritic; antiinflammatory; antiallergic; osteopathic;
RW	antisporadic; antiarteriosclerotic; antiasthmatic; immunosuppressive;
RW	neuroprotective; cardiac; cerebroprotective; cytoprotective; antidiabetic;
RW	vulner; STE20: protein kinase: STIK7: STIK3: STIK4: STIK5: STIK6: STIK7;
RW	ZC1, ZC2, ZC3, ZC4, KH52, SUDL1, SUDL3, GER2, PAK4, PAK5; antagonist;
RW	antibody; gene therapy; rheumatoid arthritis; arteriosclerosis; asthma;
RW	inflammatory bowel disease; Crohn's disease; osteoarthritis; psoriasis;
RW	rheitis; autoimmunity; organ transplantation; multiple sclerosis;
RW	myocardial infarction; cardiovascular disease; stroke; renal failure;
RW	oxidative stress-related neurodegenerative disorder; Parkinson's disease;
RW	amyotrophic lateral sclerosis; Leigh syndrome; cancer; cardiomyopathy;
RW	ischemic disorder; inflammation; diabetes mellitus; fibrosis; mitosis;
RW	mesangial disorder; growth regulation; wound healing; T cell activation;
KM	immunosuppressant.
XX	
XX	
OS	Mus sp.
XX	
PN	WO9953036-A2.
PD	21-Oct-1999.
XX	
PF	13-Apr-1999; 99WO-US08150.

[illegible]

KW	Immunosuppressant.
XX	
OS	Homo sapiens.
XX	
PN	W09953036-A2.
XX	
PD	21-OCT-1999.
XX	
PF	13-APR-1999; 99WO-US08150.
XX	
PR	14-APR-1998; 98US-0081784.
XX	
PA	(SUGEN-) SUGEN INC.
XX	
PI	Plowman G, Martinez R, Whyte D;
XX	
DR	WPI: 1999-611301/52.
XX	
PT	N-PSDB; Z40483.
XX	
PT	Novel kinase-related polypeptides used for the diagnosis and treatment
XX	of kinase-related diseases and disorders
PS	Claim 11; Page 269-274; 387pp; English.
XX	
CC	This sequence represents a novel STE20-related protein kinase. The
CC	invention relates to nucleic acid molecule encoding a kinase polypeptide
CC	selected from STEK1, STEK3, STEK4, STEK5, STEK6, STEK7, ZC1, ZC2, ZC3,
CC	ZC4, KHS2, S0U01, S0U03, GEX2, PKM4 and PKM5. The proteins are used to
CC	identify agonists and antagonists, and to raise antibodies. The
CC	polynucleotides are useful in gene therapy protocols. The polynucleotides,
CC	polypeptides, antibodies, antagonists and agonists may be used to treat
CC	diseases such as immune-related disorders and diseases (e.g. rheumatoid
CC	arthritis, atherosclerosis, chronic inflammatory bowel disease (e.g.
CC	Crohn's disease), asthma, osteoarthritis, psoriasis, atherosclerosis,
CC	rheinitis, autoimmunity, and organ transplantation, chronic inflammatory
CC	pelvic disease, multiple sclerosis, organ transplantation, myocardial
CC	infarction, cardiovascular disease, stroke, renal failure, oxidative
CC	stress-related neurodegenerative disorders (e.g. amyotrophic lateral
CC	sclerosis, Parkinson's disease and Leigh syndrome), cancer,
CC	cardiomyopathies, ischemic disorders, inflammatory disorders, diabetes
CC	mellitus, fibrotic and mesangial disorders. The proteins may also be
CC	useful for cell growth regulation (e.g. in wound healing), T cell
CC	activation, mitosis control, and as immunosuppressants.
XX	
SQ	Sequence 1239 AA;
XX	
Query Match	25.6%; Score 97.5; DB 20; Length 1239;
Best Local Similarity	39.2%; Pred. No. 0.051; Mismatches 17; Indels 11; Gaps 4
Matches 29; Conservative 17; Mismatches 17; Indels 11; Gaps 4	
QY	2 RQDPQOQYEOCQR--RCQRRTEPRHMOICQRCERYEKRRK---QQRRYEQQR- 54
DB	11: :11 11 : : 1111 1 1 :1 11 :111 : :11 11:1
DB	395 rgkrlgqkqqrllleegqrrearrqqrqr--rrgeekrllleerrrrkeeeerr 452
QY	55 --EDDEKYEERMK 66
DB	1:1::: 11 :1
DB	453 raeeekrrverege 466
XX	
RESULT 12	
Y07067	
ID	Y07067 standard; Protein; 482 AA.
XX	
AC	Y07067;
XX	
XX	
DT	02-JUL-1999 (first entry)
XX	
DE	Renal cancer associated antigen precursor sequence.
XX	
KW	Cancer associated antigen; diagnosis; research; treatment; human;
XX	breast cancer; colon cancer; gastric cancer; lung cancer; prostate cancer.
KW	

[illegible]

neuroprotective; cardiant; cerebroprotective; cytoskeletal; antidiabetic; vulnery; STE20; protein kinase; STLK2; STLK3; STLK4; STLK5; STLK6; STLK7; ZC1, ZC2, ZC3, ZC4, KHS2, SULJ1, SULJ3, GSK3, PAK4, PAK5; antagonist; antibody; gene therapy; rheumatoid arthritis; atherosclerosis; asthma; inflammatory bowel disease; Crohn's disease; osteoarthritis; psoriasis; rhinitis; autoimmunity; organ transplantation; multiple sclerosis; myocardial infarction; cardiovascular disease; stroke; renal failure; oxidative stress-related neurodegenerative disorder; Parkinson's disease; amyotrophic lateral sclerosis; Leigh syndrome; cancer; cardiomyopathy; ischemic disorder; inflammation; diabetes mellitus; fibrosis; mitosis; mesangial disorder; growth regulation; wound healing; T cell activation; immunosuppressant.

OS Homo sapiens.
 PN WO9953036-A2.
 XX 21-OCT-1999.
 PD 13-APR-1999: 99WO-US08150.
 XX 14-APR-1998: 98US-0081784.
 XX (SUGEN-) SUGEN INC.
 PA Plowman G, Martinez R, Whyte D;
 PI WPI: 1999-611301/52.
 DR N-PSDB: 240484.
 XX Novel kinase-related polypeptides used for the diagnosis and treatment of kinase-related diseases and disorders -
 PT Claim 11; Page 274-278; 387pp; English.

CC This sequence represents a novel STE20-related protein kinase. The
 CC invention relates to nucleic acid molecule encoding a kinase polypeptide
 CC selected from STLK2, STLK3, STLK4, STLK5, STLK6, STLK7, ZC1, ZC2, ZC3,
 CC ZC4, KHS2, SULJ1, SULJ3, GSK3, PAK4 and PAK5. The proteins are used to
 CC identify agonists and antagonists, and to raise antibodies. The
 CC polynucleotides are useful in gene therapy protocols. The polynucleotides,
 CC polypeptides, antibodies, antagonists and agonists may be used to treat
 CC diseases such as immune-related disorders and diseases (e.g. rheumatoid
 CC arthritis, atherosclerosis, chronic inflammatory bowel disease (e.g.
 CC Crohn's disease), asthma, osteoarthritis, psoriasis, atherosclerosis,
 CC rhinitis, autoimmunity, and organ transplantation, chronic inflammatory
 CC pelvic disease, multiple sclerosis, organ transplantation, myocardial
 CC infarction, cardiovascular disease, stroke, renal failure, oxidative
 CC stress-related neurodegenerative disorders (e.g. amyotrophic lateral
 CC sclerosis, Parkinson's disease and Leigh syndrome), cancer,
 CC cardiomyopathies, ischemic disorders, inflammatory disorders, diabetes
 CC mellitus, fibrotic and mesangial disorders. The proteins may also be
 CC useful for cell growth regulation (e.g. in wound healing), T cell
 CC activation, mitosis control, and as immunosuppressants.

XX Sequence 1297 AA;

Query Match 25.5%; Score 97; DB 20; Length 1297;
 Best Local Similarity 33.7%; Pred. No. 0.061;

Matches 31; Conservative 12; Mismatches 17; Indels 32; Gaps 5;

OY 1 NRQDP---QQQYEDCCQRCQREPRHMOICQRCR-----RYEKEX 42
 I::I I::I I::I I::I I::I I::I I::I I::I I::I I::I
 Db 327 nKseaalrtgqlq-----qgreneehkrqlaerqkrleegkeqrrrleegqrrekel 381
 OY 43 RKQQR-----YEDQOREDEEK-----YERMK 65
 I::I I::I I::I I::I I::I I::I I::I I::I I::I I::I
 Db 382 rkqgreqrthyeegmrteertraeheyk 413

RESULT 14
 Y85263

ID Y85263 standard; Protein: 1360 AA.

XX Y85263;

XX 29-JUN-2000 (first entry)

XX Human protein kinase KIAA0551 amino acid sequence.

XX Protein kinase; KIAA0551; neuropathy; neuropathic pain; inflammation;
 KW chronic pain; neurodegenerative disorder; neurotraumatic disorder;
 KW Parkinson's disease; Alzheimer's disease; ischemic disease.

XX Homo sapiens.

XX Key Location/Qualifiers

FT Misc-difference 290 /note= "Encoded by GA"

XX WO200015805-A1.

XX 23-MAR-2000.

XX 10-SEP-1999: 99WO-GB03017.

XX 10-SEP-1998: 98GB-0019779.

XX 29-MAR-1999: 99GB-0007261.

XX (SMK) SMITHKLINE BEECHAM PLC.

XX Bingham S, Case P, Lawson SN, Newton RA, Rausch OL, Reith AD;

XX Sanger GJ;

XX WPI: 2000-271443/23.

XX N-PSDB: A10669.

PT Isolated human KIAA0551 polynucleotide and polypeptide, useful for
 PT treating e.g. neuropathies, neuropathic pain, inflammatory and chronic
 PT pain and neurodegenerative conditions -
 XX Claim 2; Page 41; 48pp; English.

CC This sequence represents the human protein kinase KIAA0551 amino acid
 CC sequence. The nucleotide sequence was isolated from a human foetal brain
 CC cDNA library. The rat KIAA0551 mRNA is upregulated in dorsal root ganglia
 CC during sciatic neuropathy (a procedure accompanied by increased
 CC sensitivity to somatic pain) which indicates a role for KIAA0551 in the
 CC regulation of molecular processes associated with neuropathy and
 CC neuropathic pain. The KIAA0551 protein may be used for treating
 CC neuropathies, neuropathic pain, inflammatory and chronic pain,
 CC neurodegenerative conditions such as Parkinson's disease and Alzheimer's
 CC disease, and neurotraumatic disease or ischemic disease damage in
 CC cardiac tissue.

XX Sequence 1360 AA;

Query Match 25.2%; Score 96; DB 21; Length 1360;
 Best Local Similarity 35.4%; Pred. No. 0.08;

Matches 29; Conservative 10; Mismatches 15; Indels 28; Gaps 4;

OY 1 NRQDP---QQQYEDCCQRCQREPRHMOICQRCR-----RYEKEX 42
 I::I I::I I::I I::I I::I I::I I::I I::I I::I I::I
 Db 361 nKseaalrtgqlq-----qgreneehkrqlaerqkrleegkeqrrrleegqrrekel 415
 OY 43 RKQQR-----YEDQOREDEEK 59
 I::I I::I I::I I::I I::I I::I I::I I::I I::I I::I
 Db 416 rkqgreqrthyeegmrteer 437

RESULT 15
 Y55933
 ID Y55933 standard; Protein: 1326 AA.

